>33358 cDNA ORF 75-1046
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Figure 1A

>33358 amino acids

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SGSAACVNVLLEFGAKAQLEVHLASPIHEAVKRGHRECMEILLANNVNIDHEVPQLGTPLYVACTYQRVD
CVKKLLELGASVDHGQWLDTPLHAAARQSNVEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLL
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Figure 1B

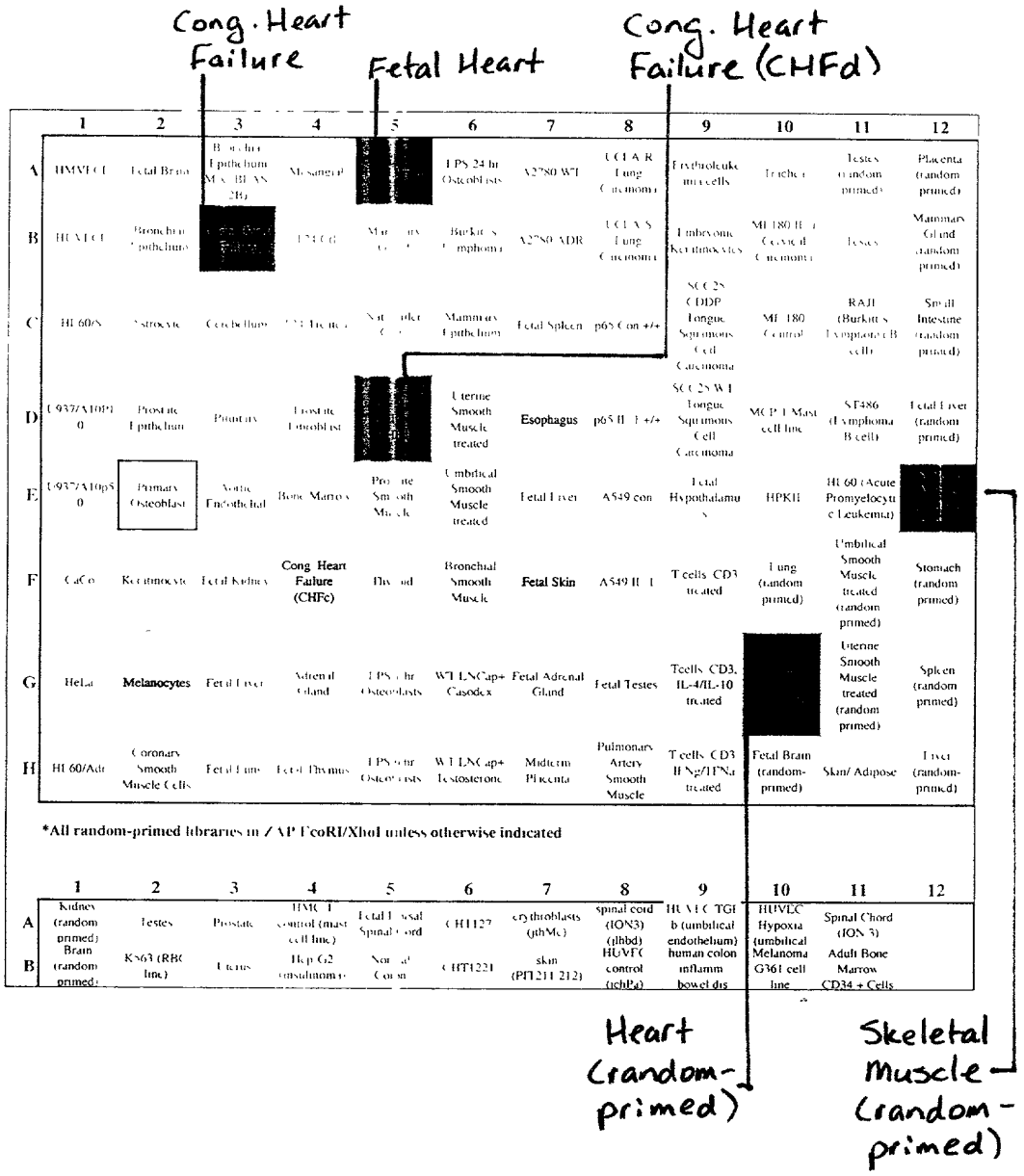

Cong. Heart Failure
Fetal Heart
Cong. Heart Failure (CHFd)
1 2 3 4 5 6 7 8 9 10 11 12
A HMVEC-I Fetal Brain Bronchial Epithelium (Mix BEAS 2B) Mesangial LPS 24 hr Osteoblasts A2780 WT LCLAR Lung Carcinoma Erythroleukemia cells Trachea Testes (random primed) Placenta (random primed)
B HUVEC-I Bronchial Epithelium LNCaP Burkitts Lymphoma A2780 ADR LCLAS Lung Carcinoma Embryonic Keratinocytes ME180 (Cervical Carcinoma) Testes Mammary Gland (random primed)
C HL60/S Astrocyte Cerebellum CD4+ T-cells Normal Colon Mammary Epithelium Fetal Spleen p65 Con +/+ SCC25 CDDP Tongue Squamous Cell Carcinoma ME 180 Control RAJI (Burkitts Lymphoma B cell) Small Intestine (random primed)
D U937/A10PI 0 Prostate Epithelium Pituitary Prostate Fibroblast Uterine Smooth Muscle treated Esophagus p65 H +/+ SCC25 WT Tongue Squamous Cell Carcinoma MCP-1 Mast cell line ST486 (Lymphoma B cell) Fetal Liver (random primed)
E U937/A10ps 0 Primary Osteoblast Aortic Endothelial Bone Marrow Prostate Smooth Muscle Umbilical Smooth Muscle treated Fetal Liver A549 con Fetal Hypothalamus HPKII HL60 (Acute Promyelocytic Leukemia)
F CaCo Keratinocyte Fetal Kidney Cong. Heart Failure (CHFc) Thyroid Bronchial Smooth Muscle Fetal Skin A549 H +/+ T cells CD3 treated Lung (random primed) Umbilical Smooth Muscle treated (random primed) Stomach (random primed)
G HeLa Melanocytes Fetal Liver Adrenal Gland LPS 4 hr Osteoclasts WT LNCap+ Casodex Fetal Adrenal Gland Fetal Testes T cells CD3, IL-4/IL-10 treated Uterine Smooth Muscle treated (random primed) Spleen (random primed)
H HL60/Adr Coronary Smooth Muscle Cells Fetal Lung Fetal Thymus LPS 4 hr Osteoclasts WT LNCap+ Testosterone Midterm Placenta Pulmonary Artery Smooth Muscle T cells CD3 IFNg/TFNa treated Fetal Brain (random-primed) Skin/ Adipose Liver (random-primed)
*All random-primed libraries in λ XP EcoRI/XhoI unless otherwise indicated
1 2 3 4 5 6 7 8 9 10 11 12
A Kidney (random primed) Testes Prostate HMC-1 control (mast cell line) Fetal Thyroid Spinal Cord CHH127 erythroblasts (jthMc) spinal cord (ION3) (jhbd) HUVEC TGFb (umbilical endothelium) human colon inflamm bowel dis HUVEC Hypoxia (umbilical Melanoma G361 cell line Spinal Chord (ION 3)
B Brain (random primed) K563 (RBC line) Uterus Hep G2 (hepatoma+ Normal Colon CHT1221 skin (PIT211 212) HUVEC control (jchPa) Adult Bone Marrow CD34 + Cells
Heart (random-primed)
Skeletal Muscle (random-primed)

Figure 2A

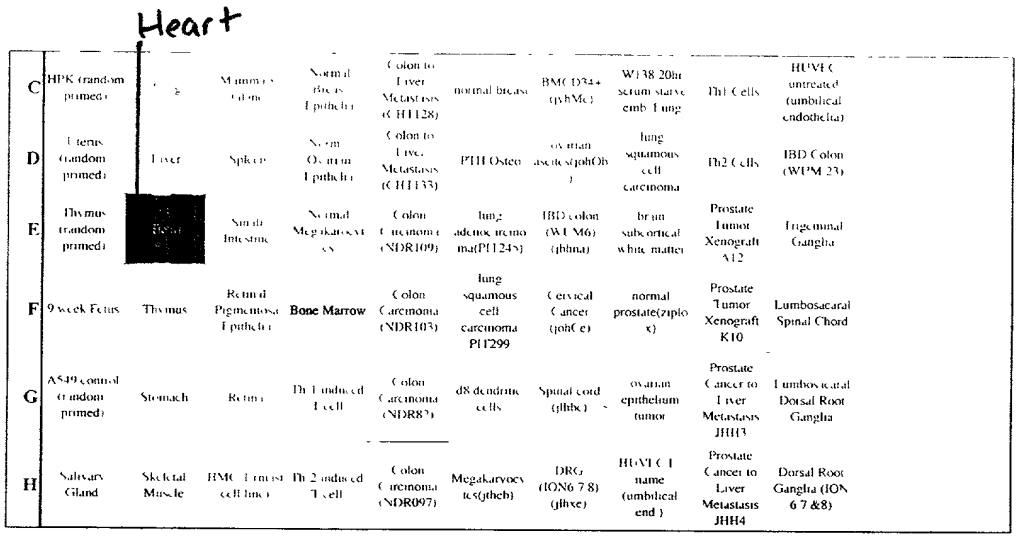
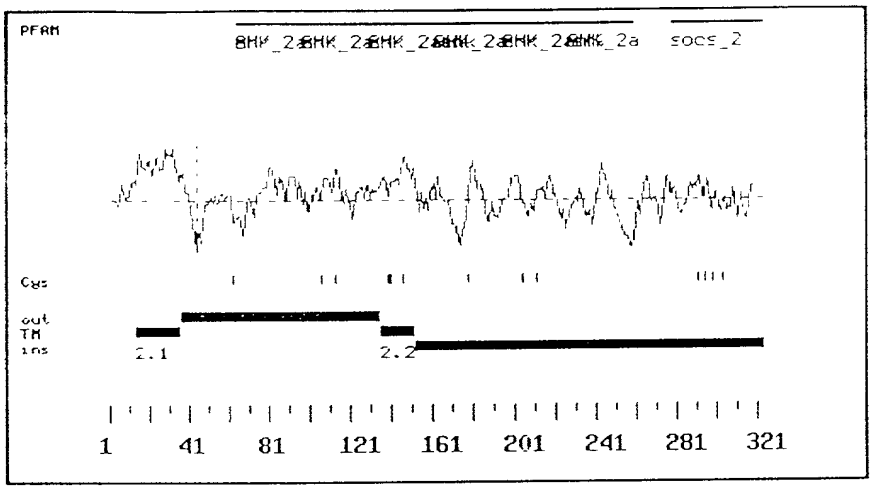

Heart
C
HPK (random primed)
Mammary Gland
Normal Breast Epithelium
Colon to Liver Metastasis (CHT128)
normal breast
BMC D3H+ (pbMc)
WI38 20hr serum starve emb Lung
Th1 Cells
HUVEC untreated (umbilical endothelial)
D
Uterus (random primed)
Liver
Spleen
Normal Ovarian Epithelium
Colon to Liver Metastasis (CHT133)
PTH Osteo
ovarian ascite(pbhOb)
lung squamous cell carcinoma
Th2 Cells
IBD Colon (WPM 23)
E
Thymus (random primed)
Small Intestine
Normal Megakaryocytes
Colon Carcinoma (NDR109)
lung adenocarcinoma(PIT243)
IBD colon (WPM6) (jbhma)
brain subcortical white matter
Prostate Tumor Xenograft A12
Trigeminal Ganglia
F
9 week Fetus
Thymus
Retinal Pigmentosa Epithelium
Bone Marrow
Colon Carcinoma (NDR103)
lung squamous cell carcinoma PIT299
Cervical Cancer (pbhCe)
normal prostate(zipl0s)
Prostate Tumor Xenograft K10
Lumbosacaral Spinal Chord
G
A549 control (random primed)
Stomach
Retina
Th1 induced T cell
Colon Carcinoma (NDR87)
d8 dendritic cells
Spinal cord (glhbc)
ovarian epithelium tumor
Prostate Cancer to Liver Metastasis JHH3
Lumbosacaral Dorsal Root Ganglia
H
Salivary Gland
Skeletal Muscle
BMC Untreated cell line
Th 2 induced T cell
Colon Carcinoma (NDR097)
Megakaryocytes(gtheb)
DRG (ION6 7 8) (glhxe)
HUVEC Untreated (umbilical end)
Prostate Cancer to Liver Metastasis JHH4
Dorsal Root Ganglia (ION 6 7 &8)

Figure 2B

Analysis of 33358 (323 aa)



>33358
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GAHVNGVTVHGATPLFNACCSGSAACVNVLLEFGAKAQLEVHLASPIHEAVKRGHRECME
ILLANNVNIDHEVPQLGTPLYVACTYQRVDCVKKLLELGASVDHGQWLDTPLHAAARQSN
VEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLLREGPPALSQLCRLCVRKCLG
RACHQAIHKLHLPEPLERFLLYQ

FIGURE 3